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1. (Twice Amended) A recombinant promoter, capable of driving expression of a transgene operably linked to the promoter, wherein the promoter comprises a nucleic acid sequence that shares at least 80% sequence identity to nucleotides 667-853 of SEQ ID NO: 17.

7. (Reiterated) A vector, comprising the recombinant promoter of claim 1.

8. (Reiterated) A host cell, comprising the vector of claim 7.

9. (Reiterated) A transgenic plant, comprising the host cell of claim 8.

10. (Reiterated) A transgene, comprising the promoter of claim 1 and at least one ORF operably linked to the promoter.

11. (Reiterated) A vector, comprising the transgene of claim 10.

12. (Reiterated) A plant cell, comprising the transgene of claim 10.

13. (Reiterated) The transgene of claim 10, wherein the ORF encodes a cationic peptide.

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14. (Twice Amended) The plant cell of claim 12, wherein the plant cell is obtained from a plant selected from the group consisting of maize, wheat, rice, millet, tobacco, sorghum, rye, barley, brassica, seaweeds, lemna, oat, soybean, cotton, legumes, rape/canola, alfalfa, flax, peanut, and clover; cucurbits, cassava, vegetables, walnuts, fruit trees, flowers, cacao; deciduous trees, conifers, turf grasses, cacao, rubber trees and members of the genus *Hevea*.

15. (Reiterated) A method for expressing at least one protein in a host cell, comprising:
introducing a transgene comprising an ORF and the recombinant promoter of claim 1 into a host cell; and
allowing the host cell to produce a protein from the ORF.

16. (Reiterated) The method of claim 15, wherein the host cell is a plant host cell.

19. (Reiterated) The recombinant promoter of claim 1, wherein the promoter is developmental-specific.

20. (Reiterated) The promoter of claim 1, wherein the promoter is induced with ethylene or a metal.

21. (Reiterated) The recombinant promoter of claim 19, wherein the promoter is expressible in gametophytic tissue.

30. (Reiterated) The plant cell of claim 14, wherein the plant cell is obtained from a tobacco plant.

31. (Reiterated) The plant cell of claim 14, wherein the plant cell is obtained from a potato plant.

32. (Reiterated) The plant cell of claim 14, wherein the plant cell is obtained from a wheat plant.

33. (Reiterated) The plant cell of claim 14, wherein the plant cell is obtained from a Douglas-fir plant.

34. (Amended) The promoter of claim 1, wherein the promoter comprises a nucleic acid sequence sharing at least 90% sequence identity to nucleotides 667-853 of SEQ ID NO: 17.

35. (Amended) The promoter of claim 1, wherein the promoter comprises [the nucleic acid sequence shown in] nucleotides 667-853 of SEQ ID NO: 17.

36. (Amended) The promoter of claim 1, wherein the promoter comprises at least 20 consecutive nucleic acid residues of a nucleic acid sequence sharing at least 80% sequence identity to nucleotides 667-853 of SEQ ID NO: 17.

37. (Amended) The promoter of claim 1, wherein the promoter comprises at least 20 consecutive nucleic acid residues of nucleotides 667-853 of SEQ ID NO: [25] 17.

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~~38~~ (Amended) The promoter of claim 1, wherein the promoter comprises at least 40 consecutive nucleic acid residues of a nucleic acid sequence sharing at least 90% sequence identity to nucleotides 667-853 of SEQ ID NO: 17.

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~~40~~ (Amended) The promoter of claim 1, wherein the promoter comprises at least 40 consecutive nucleic acid residues of nucleotides 667-853 of SEQ ID NO: [25] 17.

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~~41~~ (Amended) The promoter of claim 1, wherein the promoter comprises at least 60 consecutive nucleic acid residues of a nucleic acid sequence sharing at least 90% sequence identity to nucleotides 667-853 of SEQ ID NO: 17.

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~~42~~ (Amended) The promoter of claim 1, wherein the promoter comprises at least 60 consecutive nucleic acid residues of nucleotides 667-853 of SEQ ID NO: 17.

⁴²
~~43~~ (Amended) The promoter of claim 1, wherein the promoter comprises a nucleic acid sequence sharing at least 80% sequence identity to nucleotides 398-853 of SEQ ID NO: 17.

⁴³
~~44~~ (Amended) The promoter of claim 1, wherein the promoter comprises a nucleic acid sequence sharing at least 90% sequence identity to nucleotides 398-853 of SEQ ID NO: 17.

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~~45~~ (Amended) The promoter of claim 1, wherein the promoter comprises nucleotides 398-853 of SEQ ID NO: 17.

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~~46~~ (Amended) The promoter of claim 1, wherein the promoter comprises a nucleic acid sequence sharing at least 80% sequence identity to nucleotides 180-853 of SEQ ID NO: 17.

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~~47~~ (Amended) The promoter of claim 1, wherein the promoter comprises a nucleic acid sequence sharing at least 90% sequence identity to nucleotides 180-853 of SEQ ID NO: 17.